

## SEQUENCE LISTING

<110>	Prayaga,	Suc	hirdas	K
	Taupier	Jr,	Raymond	lJ
	Bandaru,	Raj		

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265

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<400> 24
                                                                    27
ggatccacca cctgccctc ggtgtgc
<210> 25
<211> 35
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: NOV3 Reverse
      PCR Primer Sequence
                                                                    35
ctcgaggcca gcgttctgct cctggttgag tgtgg
<210> 26
<211> 18
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: NOV3 S1 PCR
      Primer Sequence
<400> 26
                                                                    18
cgcaccattg ccagggac
<210> 27
<211> 18
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: NOV3 S2 PCR
      Primer Sequence
<400> 27
gtccctggca atggtgcg
                                                                    18
<210> 28
<211> 21
<212> DNA
<213> Artificial Sequence
<220>
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<223>	Description of Artificial Primer Sequence	Sequence:	NOV3	S3	PCR	
<400> ctggt	28 gegea attegetgge e					21
<210><211><211>	21 DNA					
<213>	Artificial Sequence					
<220> <223>	Description of Artificial Primer Sequence	Sequence:	NOV3	S4	PCR	
<400> ggcca	29 gegaa ttgegeacea g					21
<210><211><211><212><213>	18					
<220>	Description of Artificial Primer Sequence	Sequence:	NOV3	S5	PCR	
<400> cacgco	30 etctg ccaccacg					18
<210><211><211><212><213>	18					
<220> <223>	Description of Artificial Primer Sequence	Sequence:	NOV3	S6	PCR	
<400> cgtggt	31 tggca gaggcgtg					18
<210><211><211><212><213>	30					
<220> <223>	Description of Artificial Forward Oligonucleotide Pr			·V5	His	
<400> ctcgto	32 ecteg agggtaagee tateeetaad	3				30

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<210> 33
<211> 31
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: pSec-V5 His
      Reverse Oligonucleotide Primer Sequence
                                                                   31
ctcgtcgggc ccctgatcag cgggtttaaa c
<210> 34
<211> 40
<212> PRT
<213> Homo sapiens
<400> 34
Met Ala Asp Lys Pro Asp Met Gly Glu Ile Ala Ser Phe Asp Lys Ala
Lys Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Thr Leu Pro Thr Lys
                                 25
Glu Thr Ile Glu Gln Glu Lys Arg
<210> 35
<211> 10
<212> PRT
<213> Homo sapiens
<400> 35
Lys Leu Lys Lys Thr Glu Thr Gln Glu Asn
                 5
<210> 36
<211> 38
<212> PRT
<213> Homo sapiens
<400> 36
Ala Asp Lys Pro Asp Met Gly Glu Ile Ala Ser Phe Asp Lys Ala Lys
Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Thr Leu Pro Thr Lys Glu
                                  25
             20
Thr Ile Glu Gln Glu Lys
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<210> 37
<211> 40
<212> PRT
<213> Bos taurus
<400> 37
Ala Asp Lys Pro Asp Leu Gly Glu Ile Asn Ser Phe Asp Lys Ala Lys
Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Thr Leu Pro Thr Lys Glu
                                25
Thr Ile Glu Gln Glu Lys Gln Ala
<210> 38
<211> 40
<212> PRT
<213> Sus scrofa
<400> 38
Ala Asp Lys Pro Asp Met Gly Glu Ile Asn Ser Phe Asp Lys Ala Lys
Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Thr Leu Pro Thr Lys Glu
Thr Ile Glu Gln Glu Lys Gln Ala
<210> 39
<211> 40
<212> PRT
<213> Homo sapiens
<400> 39
Ser Asp Lys Pro Asp Met Ala Glu Ile Glu Lys Phe Asp Lys Ser Lys
Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Pro Leu Pro Ser Lys Glu
Thr Ile Glu Gln Glu Lys Gln Ala
<210> 40
<211> 41
<212> PRT
<213> Mus musculus
<400> 40
Met Ser Asp Lys Pro Asp Met Ala Glu Ile Glu Lys Phe Asp Lys Ser
  1 5
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Lys Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Pro Leu Pro Ser Lys 25 Glu Thr Ile Glu Gln Glu Lys Gln Ala 35 <210> 41 <211> 40 <212> PRT <213> Oryctolagus cuniculus <400> 41 Ala Asp Lys Pro Asp Met Ala Glu Ile Glu Lys Phe Asp Lys Ser Lys Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Pro Leu Pro Ser Lys Glu Thr Ile Glu Gln Glu Lys Gln Ala <210> 42 <211> 39 <212> PRT <213> Xenopus laevis <400> 42 Ser Asp Lys Pro Asp Met Ala Glu Ile Glu Lys Phe Asp Lys Ala Lys Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Pro Leu Pro Ser Lys Glu 25 Thr Ile Glu Gln Glu Lys Gln 35 <210> 43 <211> 40 <212> PRT <213> Homo sapiens <400> 43 Ser Asp Lys Pro Gly Met Ala Glu Ile Glu Lys Phe Asp Lys Ser Lys Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Pro Leu Ser Ser Lys Glu 25 Thr Ile Glu Gln Glu Arg Gln Ala

<210> 44 <211> 40

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Ser Asp Lys Pro Asn Leu Glu Glu Val Ala Ser Phe Asp Lys Thr Lys
Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Pro Leu Pro Thr Lys Glu
                                 25
Thr Ile Glu Gln Glu Lys Gln Ala
         35
<210> 45
<211> 40
<212> PRT
<213> Oncorhynchus mykiss
<400> 45
Ser Asp Lys Pro Asp Leu Ala Glu Val Ser Asn Phe Asp Lys Thr Lys
Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Pro Leu Pro Thr Lys Glu
                                 25
Thr Ile Glu Gln Glu Lys Gln Ala
         35
<210> 46
<211> 40
<212> PRT
<213> Lateolabrax japonicus
Ser Asp Lys Pro Asp Ile Ser Glu Val Thr Ser Phe Asp Lys Thr Lys
Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Pro Leu Pro Ser Lys Glu
Thr Ile Glu Gln Glu Lys Ala Ala
<210> 47
<211> 39
<212> PRT
<213> Rattus norvegicus
Met Ser Asp Lys Pro Asp Leu Ser Glu Val Glu Thr Phe Asp Lys Ser
Lys Leu Lys Lys Thr Asn Thr Glu Glu Lys Asn Thr Leu Pro Ser Lys
             20
                                 25
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<212> PRT

<213> Oncorhynchus mykiss

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<210> 48
<211> 38
<212> PRT
<213> Homo sapiens
<400> 48
Ser Asp Lys Pro Asp Leu Ser Glu Val Glu Lys Phe Asp Arg Ser Lys
Leu Lys Lys Thr Asn Thr Glu Glu Lys Asn Thr Leu Pro Ser Lys Glu
                                 25
             20
Thr Ile Gln Gln Glu Lys
         35
<210> 49
<211> 35
<212> PRT
<213> Drosophila melanogaster
<400> 49
Ile Ala Gly Ile Glu Asn Phe Asp Ala Lys Lys Leu Lys His Thr Glu
Thr Asn Glu Lys Asn Val Leu Pro Thr Lys Glu Val Ile Glu Ala Glu
                                 25
             20
Lys Gln Ala
         35
<210> 50
<211> 31
<212> PRT
<213> Drosophila melanogaster
Gly Ile Thr Ala Phe Asn Gln Asn Asn Leu Lys His Thr Glu Thr Asn
Glu Lys Asn Pro Leu Pro Asp Lys Glu Ala Ile Glu Gln Glu Lys
                                 25
             20
<210> 51
<211> 38
<212> PRT
<213> Homo sapiens
<400> 51
Ala Asp Lys Pro Asp Met Gly Glu Ile Ala Ser Phe Asp Lys Ala Lys
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Glu Thr Ile Gln Gln Glu Lys

1 5 10 15

Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Thr Leu Pro Thr Lys Glu 20 25 30

Thr Ile Glu Gln Glu Lys 35

<210> 52

<211> 991

<212> PRT

<213> Mus musculus

<400> 52

Met Ala Pro Ala Arg Ala Arg Leu Ser Pro Ala Leu Trp Val Val Thr
1 5 10 15

Ala Ala Ala Ala Thr Cys Val Ser Ala Gly Arg Gly Glu Val Asn 20 25 30

Leu Leu Asp Thr Ser Thr Ile His Gly Asp Trp Gly Trp Leu Thr Tyr
35 40 45

Pro Ala His Gly Trp Asp Ser Ile Asn Glu Val Asp Glu Ser Phe Arg
50 55 60

Pro Ile His Thr Tyr Gln Val Cys Asn Val Met Ser Pro Asn Gln Asn 65 70 75 80

Asn Trp Leu Arg Thr Asn Trp Val Pro Arg Asp Gly Ala Arg Arg Val
85 90 95

Tyr Ala Glu Ile Lys Phe Thr Leu Arg Asp Cys Asn Ser Ile Pro Gly
100 105 110

Val Leu Gly Thr Cys Lys Glu Thr Phe Asn Leu His Tyr Leu Glu Ser 115 120 125

Asp Arg Asp Leu Gly Ala Ser Thr Gln Glu Ser Gln Phe Leu Lys Ile 130 140

Asp Thr Ile Ala Ala Asp Glu Ser Phe Thr Gly Ala Asp Leu Gly Val 145 150 155 160

Arg Arg Leu Lys Leu Asn Thr Glu Val Arg Gly Val Gly Pro Leu Ser 165 170 175

Lys Arg Gly Phe Tyr Leu Ala Phe Gln Asp Ile Gly Ala Cys Leu Ala 180 185 190

Ile Leu Ser Leu Arg Ile Tyr Tyr Lys Lys Cys Pro Ala Met Val Arg 195 200 205

Asn Leu Ala Ala Phe Ser Glu Ala Val Thr Gly Ala Asp Ser Ser Ser 210 215 220

Leu 225	Val	Glu	Val	Arg	Gly 230	Gln	Cys	Val	Arg	His 235	Ser	Glu	Glu	Arg	Asp 240
Thr	Pro	Lys	Met	Tyr 245	Суѕ	Ser	Ala	Glu	Gly 250	Glu	Trp	Leu	Val	Pro 255	Ile
Gly	Lys	Cys	Val 260	Cys	Ser	Ala	Gly	Tyr 265	Glu	Glu	Arg	Arg	Asp 270	Ala	Cys
Met	Ala	Cys 275	Glu	Leu	Gly	Phe	Tyr 280	Lys	Ser	Ala	Pro	Gly 285	Asp	Gln	Leu
Cys	Ala 290	Arg	Cys	Pro	Pro	His 295	Ser	His	Ser	Ala	Thr 300	Pro	Ala	Ala	Gln
Thr 305	Cys	Arg	Cys	Asp	Leu 310	Ser	Tyr	Tyr	Arg	Ala 315	Ala	Leu	Asp	Pro	Pro 320
Ser	Ala	Ala	Cys	Thr 325	Arg	Pro	Pro	Ser	Ala 330	Pro	Val	Asn	Leu	Ile 335	Ser
Ser	Val	Asn	Gly 340	Thr	Ser	Val	Thr	Leu 345	Glu	Trp	Ala	Pro	Pro 350	Leu	Asp
Pro	Gly	Gly 355	Arg	Ser	Asp	Ile	Thr 360	Tyr	Asn	Ala	Val	Cys 365	Arg	Arg	Cys
Pro	Trp 370	Ala	Leu	Ser	His	Cys 375	Glu	Ala	Cys	Gly	Ser 380	Gly	Thr	Arg	Phe
Val 385	Pro	Gln	Gln	Thr	Ser 390	Leu	Ala	Gln	Ala	Ser 395	Leu	Leu	Val	Ala	Asn 400
Leu	Leu	Ala	His	Met 405	Asn	Tyr	Ser	Phe	Trp 410	Ile	Glu	Ala	Val	Asn 415	Gly
Val	Ser	Asn	Leu 420	Ser	Pro	Glu	Pro	Arg 425	Ser	Ala	Ala	Val	Val 430	Asn	Ile
Thr	Thr	Asn 435	Gln	Ala	Ala	Pro	Ser 440	Gln	Val	Val	Val	Ile 445	Arg	Gln	Glu
Arg	Ala 450	Gly	Gln	Thr	Ser	Val 455	Ser	Leu	Leu	Trp	Gln 460	Glu	Pro	Glu	Gln
Pro 465	Asn	Gly	Ile	Ile	Leu 470	Glu	Tyr	Glu	Ile	Lys 475	Tyr	Tyr	Glu	Lys	Asp 480
Lys	Glu	Met	Gln	Ser 485	Tyr	Ser	Thr	Leu	Lys 490	Ala	Val	Thr	Thr	Arg 495	Ala
Thr	Val	Ser	Gly 500	Leu	Lys	Pro	Gly	Thr 505	Arg	Tyr	Val	Phe	Gln 510	Val	Arg
Ala	Arg	Thr 515	Ser	Ala	Gly	Cys	Gly 520	Arg	Phe	Ser	Gln	Ala 525	Met	Glu	Val

Glu Thr Gly Lys Pro Arg Pro Arg Tyr Asp Thr Arg Thr Ile Val Trp 535 Ile Cys Leu Thr Leu Ile Thr Gly Leu Val Val Leu Leu Leu Leu Leu 555 Ile Cys Lys Lys Arg His Cys Gly Tyr Ser Lys Ala Phe Gln Asp Ser Asp Glu Glu Lys Met His Tyr Gln Asn Gly Gln Ala Pro Pro Pro Val 585 Phe Leu Pro Leu Asn His Pro Pro Gly Lys Phe Pro Glu Thr Gln Phe Ser Ala Glu Pro His Thr Tyr Glu Glu Pro Gly Arg Ala Gly Arg Ser Phe Thr Arg Glu Ile Glu Ala Ser Arg Ile His Ile Glu Lys Ile Ile Gly Ser Gly Glu Ser Gly Glu Val Cys Tyr Gly Arg Leu Gln Val Pro Gly Gln Arg Asp Val Pro Val Ala Ile Lys Ala Leu Lys Ala Gly Tyr 665 Thr Glu Arg Gln Arg Gln Asp Phe Leu Ser Glu Ala Ala Ile Met Gly 680 Gln Phe Asp His Pro Asn Ile Ile Arg Leu Glu Gly Val Val Thr Arg 695 Gly Arg Leu Ala Met Ile Val Thr Glu Tyr Met Glu Asn Gly Ser Leu 715 Asp Ala Phe Leu Arg Thr His Asp Gly Gln Phe Thr Ile Val Gln Leu Val Gly Met Leu Arg Gly Val Gly Ala Gly Met Arg Tyr Leu Ser Asp 745 Leu Gly Tyr Ile His Arg Asp Leu Ala Ala Arg Asn Val Leu Val Asp Gly Arg Leu Val Cys Lys Val Ser Asp Phe Gly Leu Ser Arg Ala Leu 775 Glu Asp Asp Pro Glu Ala Ala Tyr Thr Thr Ala Gly Gly Lys Ile Pro 795 Ile Arg Trp Thr Ala Pro Glu Ala Ile Ala Phe Arg Thr Phe Ser Ser Ala Ser Asp Val Trp Ser Phe Gly Val Val Met Trp Glu Val Leu Ala 820 825

Tyr Gly Glu Arg Pro Tyr Trp Asn Met Thr Asn Gln Asp Val Ile Ser 835 840 845

Ser Val Glu Glu Gly Tyr Arg Leu Pro Ala Pro Met Gly Cys Pro Arg 850 855 860

Ala Leu His Gln Leu Met Leu Asp Cys Trp His Lys Asp Arg Ala Gln 865 870 875 880

Arg Pro Arg Phe Ala His Val Val Ser Val Leu Asp Ala Leu Val His 885 890 895

Ser Pro Glu Ser Leu Arg Ala Thr Ala Thr Val Ser Arg Cys Pro Pro 900 905 910

Pro Ala Phe Ala Arg Ser Cys Phe Asp Leu Arg Ala Gly Gly Ser Gly 915 920 925

Asn Gly Asp Leu Thr Val Gly Asp Trp Leu Asp Ser Ile Arg Met Gly 930 935 940

Arg Tyr Arg Asp His Phe Ala Ala Gly Gly Tyr Ser Ser Leu Gly Met 945 950 955 960

Val Leu Arg Met Asn Ala Gln Asp Val Arg Ala Leu Gly Ile Thr Leu
965 970 975

Met Gly His Gln Lys Lys Ile Leu Gly Ser Ile Gln Thr Met Arg 980 985 990

<210> 53

<211> 992

<212> PRT

<213> Homo sapiens

<400> 53

Met Ala Pro Ala Arg Gly Arg Leu Pro Pro Ala Leu Trp Val Val Thr 1 5 10 15

Ala Ala Ala Ala Ala Thr Cys Val Ser Ala Ala Arg Gly Glu Val 20 25 30

Asn Leu Leu Asp Thr Ser Thr Ile His Gly Asp Trp Gly Trp Leu Thr 35 40 45

Tyr Pro Ala His Gly Trp Asp Ser Ile Asn Glu Val Asp Glu Ser Phe 50 55 60

Gln Pro Ile His Thr Tyr Gln Val Cys Asn Val Met Ser Pro Asn Gln 65 70 75 80

Asn Asn Trp Leu Arg Thr Ser Trp Val Pro Arg Asp Gly Ala Arg Arg 85 90 95

Val Tyr Ala Glu Ile Lys Phe Thr Leu Arg Asp Cys Asn Ser Met Pro 100 105 110 Gly Val Leu Gly Thr Cys Lys Glu Thr Phe Asn Leu Tyr Tyr Leu Glu Ser Asp Arg Asp Leu Gly Ala Ser Thr Gln Glu Ser Gln Phe Leu Lys 135 Ile Asp Thr Ile Ala Ala Asp Glu Ser Phe Thr Gly Ala Asp Leu Gly Val Arg Arg Leu Lys Leu Asn Thr Glu Val Arg Ser Val Gly Pro Leu 165 Ser Lys Arg Gly Phe Tyr Leu Ala Phe Gln Asp Ile Gly Ala Cys Leu 185 Ala Ile Leu Ser Leu Arg Ile Tyr Tyr Lys Lys Cys Pro Ala Met Val 200 Arg Asn Leu Ala Ala Phe Ser Glu Ala Val Thr Gly Ala Asp Ser Ser 220 Ser Leu Val Glu Val Arg Gly Gln Cys Val Arg His Ser Glu Glu Arg 225 Asp Thr Pro Lys Met Tyr Cys Ser Ala Glu Gly Glu Trp Leu Val Pro 250 Ile Gly Lys Cys Val Cys Ser Ala Gly Tyr Glu Glu Arg Arg Asp Ala Cys Val Ala Cys Glu Leu Gly Phe Tyr Lys Ser Ala Pro Gly Asp Gln 280 Leu Cys Ala Arg Cys Pro Pro His Ser His Ser Ala Ala Pro Ala Ala 295 Gln Ala Cys His Cys Asp Leu Ser Tyr Tyr Arg Ala Ala Leu Asp Pro 315 Pro Ser Ser Ala Cys Thr Arg Pro Pro Ser Ala Pro Val Asn Leu Ile Ser Ser Val Asn Gly Thr Ser Val Thr Leu Glu Trp Ala Pro Pro Leu 345 Asp Pro Gly Gly Arg Ser Asp Ile Thr Tyr Asn Ala Val Cys Arg Arg 360 Cys Pro Trp Ala Leu Ser Arg Cys Glu Ala Cys Gly Ser Gly Thr Arg 375 Phe Val Pro Gln Gln Thr Ser Leu Val Gln Ala Ser Leu Leu Val Ala 395 Asn Leu Leu Ala His Met Asn Tyr Ser Phe Trp Ile Glu Ala Val Asn 405 410

Gly Val Ser Asp Leu Ser Pro Glu Pro Arg Arg Ala Ala Val Val Asn Ile Thr Thr Asn Gln Ala Ala Pro Ser Gln Val Val Val Ile Arg Gln Glu Arg Ala Gly Gln Thr Ser Val Ser Leu Leu Trp Gln Glu Pro Glu 455 Gln Pro Asn Gly Ile Ile Leu Glu Tyr Glu Ile Lys Tyr Tyr Glu Lys Asp Lys Glu Met Gln Ser Tyr Ser Thr Leu Lys Ala Val Thr Thr Arg Ala Thr Val Ser Gly Leu Lys Pro Gly Thr Arg Tyr Val Phe Gln Val Arg Ala Arg Thr Ser Ala Gly Cys Gly Arg Phe Ser Gln Ala Met Glu Val Glu Thr Gly Lys Pro Arg Pro Arg Tyr Asp Thr Arg Thr Ile Val 535 Trp Ile Cys Leu Thr Leu Ile Thr Gly Leu Val Val Leu Leu Leu 550 555 Leu Ile Cys Lys Lys Arg His Cys Gly Tyr Ser Lys Ala Phe Gln Asp 570 Ser Asp Glu Glu Lys Met His Tyr Gln Asn Gly Gln Ala Pro Pro Pro Val Phe Leu Pro Leu His His Pro Pro Gly Lys Leu Pro Glu Pro Gln 595 Phe Tyr Ala Glu Pro His Thr Tyr Glu Glu Pro Gly Arg Ala Gly Arg 615 Ser Phe Thr Arg Glu Ile Glu Ala Ser Arg Ile His Ile Glu Lys Ile 625 Ile Gly Ser Gly Asp Ser Gly Glu Val Cys Tyr Gly Arg Leu Arg Val Pro Gly Gln Arg Asp Val Pro Val Ala Ile Lys Ala Leu Lys Ala Gly Tyr Thr Glu Arg Gln Arg Arg Asp Phe Leu Ser Glu Ala Ser Ile Met Gly Gln Phe Asp His Pro Asn Ile Ile Arg Leu Glu Gly Val Val Thr 695 690 Arg Gly Arg Leu Ala Met Ile Val Thr Glu Tyr Met Glu Asn Gly Ser 705 710 715

- Leu Asp Thr Phe Leu Arg Thr His Asp Gly Gln Phe Thr Ile Met Gln 725 730 735
- Leu Val Gly Met Leu Arg Gly Val Gly Ala Gly Met Arg Tyr Leu Ser
  740 745 750
- Asp Leu Gly Tyr Val His Arg Asp Leu Ala Ala Arg Asn Val Leu Val 755 760 765
- Asp Ser Asn Leu Val Cys Lys Val Ser Asp Phe Gly Leu Ser Arg Val 770 780
- Leu Glu Asp Asp Pro Asp Ala Ala Tyr Thr Thr Thr Gly Gly Lys Ile
  785 790 795 800
- Pro Ile Arg Trp Thr Ala Pro Glu Ala Ile Ala Phe Arg Thr Phe Ser 805 810 815
- Ser Ala Ser Asp Val Trp Ser Phe Gly Val Val Met Trp Glu Val Leu 820 825 830
- Ala Tyr Gly Glu Arg Pro Tyr Trp Asn Met Thr Asn Arg Asp Val Ile 835 840 845
- Ser Ser Val Glu Glu Gly Tyr Arg Leu Pro Ala Pro Met Gly Cys Pro 850 860
- His Ala Leu His Gln Leu Met Leu Asp Cys Trp His Lys Asp Arg Ala 865 870 875 880
- Gln Arg Pro Arg Phe Ser Gln Ile Val Ser Val Leu Asp Ala Leu Ile 885 890 895
- Pro Pro Ala Phe Val Arg Ser Cys Phe Asp Leu Arg Gly Gly Ser Gly 915 920 925
- Gly Gly Gly Leu Thr Val Gly Asp Trp Leu Asp Ser Ile Arg Met 930 935 940
- Gly Arg Tyr Arg Asp His Phe Ala Ala Gly Gly Tyr Ser Ser Leu Gly 945 950 955 960
- Met Val Leu Arg Met Asn Ala Gln Asp Val Arg Ala Leu Gly Ile Thr 965 970 975
- Leu Met Gly His Gln Lys Lys Ile Leu Gly Ser Ile Gln Thr Met Arg 980 985 990

- <211> 450
- <212> PRT
- <213> Mus musculus
- <400> 54
- Met Ala Pro Ala Arg Ala Arg Leu Ser Pro Ala Leu Trp Val Val Thr
  1 5 10 15
- Ala Ala Ala Ala Thr Cys Val Ser Ala Gly Arg Gly Glu Val Asn 20 25 30
- Leu Leu Asp Thr Ser Thr Ile His Gly Asp Trp Gly Trp Leu Thr Tyr 35 40 45
- Pro Ala His Gly Trp Asp Ser Ile Asn Glu Val Asp Glu Ser Phe Arg
  50 60
- Pro Ile His Thr Tyr Gln Val Cys Asn Val Met Ser Pro Asn Gln Asn 65 70 75 80
- Asn Trp Leu Arg Thr Asn Trp Val Pro Arg Asp Gly Ala Arg Arg Val
- Tyr Ala Glu Ile Lys Phe Thr Leu Arg Asp Cys Asn Ser Ile Pro Gly
  100 105 110
- Val Leu Gly Thr Cys Lys Glu Thr Phe Asn Leu His Tyr Leu Glu Ser 115 120 125
- Asp Arg Asp Leu Gly Ala Ser Thr Gln Glu Ser Gln Phe Leu Lys Ile 130 135 140
- Asp Thr Ile Ala Ala Asp Glu Ser Phe Thr Gly Ala Asp Leu Gly Val 145 150 155 160
- Arg Arg Leu Lys Leu Asn Thr Glu Val Arg Gly Val Gly Pro Leu Ser 165 170 175
- Lys Arg Gly Phe Tyr Leu Ala Phe Gln Asp Ile Gly Ala Cys Leu Ala 180 185 190
- Ile Leu Ser Leu Arg Ile Tyr Tyr Lys Lys Cys Pro Ala Met Val Arg 195 200 205
- Asn Leu Ala Ala Phe Ser Glu Ala Val Thr Gly Ala Asp Ser Ser Ser 210 215 220
- Leu Val Glu Val Arg Gly Gln Cys Val Arg His Ser Glu Glu Arg Asp 225 230 235 240
- Thr Pro Lys Met Tyr Cys Ser Ala Glu Gly Glu Trp Leu Val Pro Ile 245 250 255
- Gly Lys Cys Val Cys Ser Ala Gly Tyr Glu Glu Arg Arg Asp Ala Cys 260 265 270
- Met Ala Cys Glu Leu Gly Phe Tyr Lys Ser Ala Pro Gly Asp Gln Leu

275 280 285

Cys Ala Arg Cys Pro Pro His Ser His Ser Ala Thr Pro Ala Ala Gln 290 295 300

Thr Cys Arg Cys Asp Leu Ser Tyr Tyr Arg Ala Ala Leu Asp Pro Pro 305 310 315 320

Ser Ala Ala Cys Thr Arg Pro Pro Ser Ala Pro Val Asn Leu Ile Ser 325 330 335

Ser Val Asn Gly Thr Ser Val Thr Leu Glu Trp Ala Pro Pro Leu Asp 340 345 350

Pro Gly Gly Arg Ser Asp Ile Thr Tyr Asn Ala Val Cys Arg Arg Cys 355 360 365

Pro Trp Ala Leu Ser His Cys Glu Ala Cys Gly Ser Gly Thr Arg Phe 370 380

Val Pro Gln Gln Thr Ser Leu Ala Gln Ala Ser Leu Leu Val Ala Asn 385 390 395 400

Leu Leu Ala His Met Asn Tyr Ser Phe Trp Ile Glu Ala Val Asn Gly
405 410 415

Val Ser Asn Leu Ser Pro Glu Pro Arg Ser Ala Ala Val Val Asn Ile 420 425 430

Thr Thr Asn Gln Ala Ala Pro Ser Gln Val Val Val Ile Arg Gln Glu
435 440 445

Arg Ala 450

<210> 55

<211> 480

<212> PRT

<213> Homo sapiens

<400> 55

Met Arg Gly Ser Gly Pro Arg Gly Ala Gly His Arg Arg Pro Pro Ser 1 5 10 15

Gly Gly Gly Asp Thr Pro Ile Thr Pro Ala Ser Leu Ala Gly Cys Tyr 20 25 30

Ser Ala Pro Arg Ala Pro Leu Trp Thr Cys Leu Leu Cys Ala 35 40 45

Ala Leu Arg Thr Leu Leu Ala Ser Pro Ser Asn Glu Val Asn Leu Leu 50 60

Asp Ser Arg Thr Val Met Gly Asp Leu Gly Trp Ile Ala Phe Pro Lys 65 70 75 80

Asn Gly Trp Glu Glu Ile Gly Glu Val Asp Glu Asn Tyr Ala Pro Ile His Thr Tyr Gln Val Cys Lys Val Met Glu Gln Asn Gln Asn Asn Trp Leu Leu Thr Ser Trp Ile Ser Asn Glu Gly Ala Ser Arg Ile Phe Ile 120 Glu Leu Lys Phe Thr Leu Arg Asp Cys Asn Ser Leu Pro Gly Gly Leu 135 Gly Thr Cys Lys Glu Thr Phe Asn Met Tyr Tyr Phe Glu Ser Asp Asp Gln Asn Gly Arg Asn Ile Lys Glu Asn Gln Tyr Ile Lys Ile Asp Thr Ile Ala Ala Asp Glu Ser Phe Thr Glu Leu Asp Leu Gly Asp Arg Val 180 185 Met Lys Leu Asn Thr Glu Val Arg Asp Val Gly Pro Leu Ser Lys Lys 200 Gly Phe Tyr Leu Ala Phe Gln Asp Val Gly Ala Cys Ile Ala Leu Val 215 Ser Val Arg Val Tyr Tyr Lys Lys Cys Pro Ser Val Val Arg His Leu 235 Ala Val Phe Pro Asp Thr Ile Thr Gly Ala Asp Ser Ser Gln Leu Leu 250 Glu Val Ser Gly Ser Cys Val Asn His Ser Val Thr Asp Glu Pro Pro 265 Lys Met His Cys Ser Ala Glu Gly Glu Trp Leu Val Pro Ile Gly Lys Cys Met Cys Lys Ala Gly Tyr Glu Glu Lys Asn Gly Thr Cys Gln Val 295 Cys Arg Pro Gly Phe Phe Lys Ala Ser Pro His Ile Gln Ser Cys Gly 305 Lys Cys Pro Pro His Ser Tyr Thr His Glu Glu Ala Ser Thr Ser Cys 330 Val Cys Glu Lys Asp Tyr Phe Arg Arg Glu Ser Asp Pro Pro Thr Met Ala Cys Thr Arg Pro Pro Ser Ala Pro Arg Asn Ala Ile Ser Asn Val Asn Glu Thr Ser Val Phe Leu Glu Trp Ile Pro Pro Ala Asp Thr Gly 370 375 380

Gly Arg Lys Asp Val Ser Tyr Tyr Ile Ala Cys Lys Lys Cys Asn Ser 385 390 395 400

His Ala Gly Val Cys Glu Glu Cys Gly Gly His Val Arg Tyr Leu Pro
405 410 415

Arg Gln Ser Gly Leu Lys Asn Thr Ser Val Met Met Val Asp Leu Leu 420 425 430

Ala His Thr Asn Tyr Thr Phe Glu Ile Glu Ala Val Asn Gly Val Ser 435 440 445

Asp Leu Ser Pro Gly Ala Arg Gln Tyr Val Ser Val Asn Val Thr Thr 450 455 460

Asn Gln Ala Ala Pro Ser Pro Val Thr Asn Val Lys Lys Gly Lys Ile 465 470 475 480

<210> 56

<211> 456

<212> PRT

<213> Gallus gallus

<400> 56

Met Gly Leu Arg Gly Gly Gly Gly Arg Ala Gly Gly Pro Ala Pro Gly
1 5 10 15

Trp Thr Cys Leu Leu Cys Ala Ala Leu Arg Ser Leu Leu Ala Ser
20 25 30

Pro Gly Ser Glu Val Asn Leu Leu Asp Ser Arg Thr Val Met Gly Asp 35 40 45

Leu Gly Trp Ile Ala Tyr Pro Lys Asn Gly Trp Glu Glu Ile Gly Glu
50 60

Val Asp Glu Asn Tyr Ala Pro Ile His Thr Tyr Gln Val Cys Lys Val 65 70 75 80

Met Glu Gln Asn Gln Asn Asn Trp Leu Leu Thr Ser Trp Ile Ser Asn 85 90 95

Glu Gly Arg Pro Ala Ser Ser Phe Glu Leu Lys Phe Thr Leu Arg Asp 100 105 110

Cys Asn Ser Leu Pro Gly Gly Leu Gly Thr Cys Lys Glu Thr Phe Asn 115 120 125

Met Tyr Tyr Phe Glu Ser Asp Asp Glu Asp Gly Arg Asn Ile Arg Glu 130 135 140

Asn Gln Tyr Ile Lys Ile Asp Thr Ile Ala Ala Asp Glu Ser Phe Thr 145 150 155 160

Glu Leu Asp Leu Gly Asp Arg Val Met Lys Leu Asn Thr Glu Val Arg 170 165 Asp Val Gly Pro Leu Thr Lys Lys Gly Phe Tyr Leu Ala Phe Gln Asp 185 Val Gly Ala Cys Ile Ala Leu Val Ser Val Arg Val Tyr Tyr Lys Lys Cys Pro Ser Val Ile Arg Asn Leu Ala Arg Phe Pro Asp Thr Ile Thr Gly Ala Asp Ser Ser Gln Leu Leu Glu Val Ser Gly Val Cys Val Asn His Ser Val Thr Asp Glu Ala Pro Lys Met His Cys Ser Ala Glu Gly Glu Trp Leu Val Pro Ile Gly Lys Cys Leu Cys Lys Ala Gly Tyr Glu Glu Lys Asn Asn Thr Cys Gln Val Cys Arg Pro Gly Phe Phe Lys Ala Ser Pro His Ser Pro Ser Cys Ser Lys Cys Pro Pro His Ser Tyr Thr 295 Leu Asp Glu Ala Ser Thr Ser Cys Leu Cys Glu Glu His Tyr Phe Arg 310 Arg Glu Ser Asp Pro Pro Thr Met Ala Cys Thr Arg Pro Pro Ser Ala 325 330 Pro Arq Ser Ala Ile Ser Asn Val Asn Glu Thr Ser Val Phe Leu Glu 345 Trp Ile Pro Pro Ala Asp Thr Gly Gly Arg Lys Asp Val Ser Tyr Tyr Ile Ala Cys Lys Lys Cys Asn Ser His Ser Gly Leu Cys Glu Ala Cys Gly Ser His Val Arg Tyr Leu Pro Gln Gln Thr Gly Leu Lys Asn Thr 390 395 Ser Val Met Met Val Asp Leu Leu Ala His Thr Asn Tyr Thr Phe Glu 405 Ile Glu Ala Val Asn Gly Val Ser Asp Gln Asn Pro Gly Ala Arg Gln 425 Phe Val Ser Val Asn Val Thr Thr Asn Gln Ala Ala Pro Ser Pro Val Ser Ser Val Lys Lys Gly Lys Ile

455

- <210> 57 <211> 649
- <211> 645
- <212> PRT
- <213> Homo sapiens
- <400> 57
- Met Ile Ser Ala Ala Trp Ser Ile Phe Leu Ile Gly Thr Lys Ile Gly

  1 10 15
- Leu Phe Leu Gln Val Ala Pro Leu Ser Val Met Ala Lys Ser Cys Pro 20 25 30
- Ser Val Cys Arg Cys Asp Ala Gly Phe Ile Tyr Cys Asn Asp Arg Phe 35 40 45
- Leu Thr Ser Ile Pro Thr Gly Ile Pro Glu Asp Ala Thr Thr Leu Tyr
  50 55 60
- Leu Gln Asn Asn Gln Ile Asn Asn Ala Gly Ile Pro Ser Asp Leu Lys
  65 70 75 80
- Asn Leu Leu Lys Val Glu Arg Ile Tyr Leu Tyr His Asn Ser Leu Asp 85 90 95
- Glu Phe Pro Thr Asn Leu Pro Lys Tyr Val Lys Glu Leu His Leu Gln
  100 105 110
- Glu Asn Asn Ile Arg Thr Ile Thr Tyr Asp Ser Leu Ser Lys Ile Pro 115 120 125
- Tyr Leu Glu Glu Leu His Leu Asp Asp Asn Ser Val Ser Ala Val Ser 130 135 140
- Ile Glu Glu Gly Ala Phe Arg Asp Ser Asn Tyr Leu Arg Leu Leu Phe 145 150 155 160
- Leu Ser Arg Asn His Leu Ser Thr Ile Pro Trp Gly Leu Pro Arg Thr
  165 170 175
- Ile Glu Glu Leu Arg Leu Asp Asp Asn Arg Ile Ser Thr Ile Ser Ser 180 185 190
- Pro Ser Leu Gln Gly Leu Thr Ser Leu Lys Arg Leu Val Leu Asp Gly 195 200 205
- Asn Leu Leu Asn Asn His Gly Leu Gly Asp Lys Val Phe Phe Asn Leu 210 215 220
- Val Asn Leu Thr Glu Leu Ser Leu Val Arg Asn Ser Leu Thr Ala Ala 225 230 235 240
- Pro Val Asn Leu Pro Gly Thr Asn Leu Arg Lys Leu Tyr Leu Gln Asp 245 250 255
- Asn His Ile Asn Arg Val Pro Pro Asn Ala Phe Ser Tyr Leu Arg Gln

260 265 270

Leu Tyr Arg Leu Asp Met Ser Asn Asn Asn Leu Ser Asn Leu Pro Gln 280 Gly Ile Phe Asp Asp Leu Asp Asn Ile Thr Gln Leu Ile Leu Arg Asn 295 Asn Pro Trp Tyr Cys Gly Cys Lys Met Lys Trp Val Arg Asp Trp Leu Gln Ser Leu Pro Val Lys Val Asn Val Arg Gly Leu Met Cys Gln Ala 330 Pro Glu Lys Val Arg Gly Met Ala Ile Lys Asp Leu Asn Ala Glu Leu 345 Phe Asp Cys Lys Asp Ser Gly Ile Val Ser Thr Ile Gln Ile Thr Thr 360 Ala Ile Pro Asn Thr Val Tyr Pro Ala Gln Gly Gln Trp Pro Ala Pro 375 Val Thr Lys Gln Pro Asp Ile Lys Asn Pro Lys Leu Thr Lys Asp His 395 390 Gln Thr Thr Gly Ser Pro Ser Arg Lys Thr Ile Thr Ile Thr Val Lys Ser Val Thr Ser Asp Thr Ile His Ile Ser Trp Lys Leu Ala Leu Pro 425 Met Thr Ala Leu Arg Leu Ser Trp Leu Lys Leu Gly His Ser Pro Ala Phe Gly Ser Ile Thr Glu Thr Ile Val Thr Gly Glu Arg Ser Glu Tyr Leu Val Thr Ala Leu Glu Pro Asp Ser Pro Tyr Lys Val Cys Met Val 465 Pro Met Glu Thr Ser Asn Leu Tyr Leu Phe Asp Glu Thr Pro Val Cys 485 Ile Glu Thr Glu Thr Ala Pro Leu Arg Met Tyr Asn Pro Thr Thr 505 500 Leu Asn Arg Glu Gln Glu Lys Glu Pro Tyr Lys Asn Pro Asn Leu Pro Leu Ala Ala Ile Ile Gly Gly Ala Val Ala Leu Val Thr Ile Ala Leu Leu Ala Leu Val Cys Trp Tyr Val His Arg Asn Gly Ser Leu Phe Ser

Arg Asn Cys Ala Tyr Ser Lys Gly Arg Arg Arg Lys Asp Asp Tyr Ala

565 570 575

Glu Ala Gly Thr Lys Lys Asp Asn Ser Ile Leu Glu Ile Arg Glu Thr 580 585 590

Ser Phe Gln Met Leu Pro Ile Ser Asn Glu Pro Ile Ser Lys Glu Glu 595 600 605

Phe Val Ile His Thr Ile Phe Pro Pro Asn Gly Met Asn Leu Tyr Lys 610 615 620

Asn Asn His Ser Glu Ser Ser Ser Asn Arg Ser Tyr Arg Asp Ser Gly 625 630 635

Ile Pro Asp Ser Asp His Ser His Ser 645

<210> 58

<211> 660

<212> PRT

<213> Homo sapiens

<400> 58

Met Gly Leu Gln Thr Thr Lys Trp Pro Ser His Gly Ala Phe Phe Leu 1 5 10 15

Lys Ser Trp Leu Ile Ile Ser Leu Gly Leu Tyr Ser Gln Val Ser Lys 20 25 30

Leu Leu Ala Cys Pro Ser Val Cys Arg Cys Asp Arg Asn Phe Val Tyr
35 40 45

Cys Asn Glu Arg Ser Leu Thr Ser Val Pro Leu Gly Ile Pro Glu Gly 50 55 60

Val Thr Val Leu Tyr Leu His Asn Asn Gln Ile Asn Asn Ala Gly Phe
65 70 75 80

Pro Ala Glu Leu His Asn Val Gln Ser Val His Thr Val Tyr Leu Tyr
85 90 95

Gly Asn Gln Leu Asp Glu Phe Pro Met Asn Leu Pro Lys Asn Val Arg 100 105 110

Val Leu His Leu Gln Glu Asn Asn Ile Gln Thr Ile Ser Arg Ala Ala 115 120 125

Leu Ala Gln Leu Leu Lys Leu Glu Glu Leu His Leu Asp Asp Asn Ser 130 135 140

Ile Ser Thr Val Gly Val Glu Asp Gly Ala Phe Arg Glu Ala Ile Ser 145 150 155 160

Leu Lys Leu Leu Phe Leu Ser Lys Asn His Leu Ser Ser Val Pro Val
165 170 175

Gly Leu Pro Val Asp Leu Gln Glu Leu Arg Val Asp Glu Asn Arg Ile Ala Val Ile Ser Asp Met Ala Phe Gln Asn Leu Thr Ser Leu Glu Arg Leu Ile Val Asp Gly Asn Leu Leu Thr Asn Lys Gly Ile Ala Glu Gly 215 Thr Phe Ser His Leu Thr Lys Leu Lys Glu Phe Ser Ile Val Arg Asn 235 Ser Leu Ser His Pro Pro Pro Asp Leu Pro Gly Thr His Leu Ile Arg Leu Tyr Leu Gln Asp Asn Gln Ile Asn His Ile Pro Leu Thr Ala Phe 265 Ser Asn Leu Arg Lys Leu Glu Arg Leu Asp Ile Ser Asn Asn Gln Leu 280 Arg Met Leu Thr Gln Gly Val Phe Asp Asn Leu Ser Asn Leu Lys Gln 295 Leu Thr Ala Arg Asn Asn Pro Trp Phe Cys Asp Cys Ser Ile Lys Trp 310 315 Val Thr Glu Trp Leu Lys Tyr Ile Pro Ser Ser Leu Asn Val Arg Gly 330 Phe Met Cys Gln Gly Pro Glu Gln Val Arg Gly Met Ala Val Arg Glu Leu Asn Met Asn Leu Leu Ser Cys Pro Thr Thr Pro Gly Leu Pro 360 Leu Phe Thr Pro Ala Pro Ser Thr Ala Ser Pro Thr Thr Gln Pro Pro 370 Thr Leu Ser Ile Pro Asn Pro Ser Arg Ser Tyr Thr Pro Pro Thr Pro 390 Thr Thr Ser Lys Leu Pro Thr Ile Pro Asp Trp Asp Gly Arg Glu Arg 405 Val Thr Pro Pro Ile Ser Glu Arg Ile Gln Leu Ser Ile His Phe Val 425 Asn Asp Thr Ser Ile Gln Val Ser Trp Leu Ser Leu Phe Thr Val Met Ala Tyr Lys Leu Thr Trp Val Lys Met Gly His Ser Leu Val Gly Gly 455 Ile Val Gln Glu Arg Ile Val Ser Gly Glu Lys Gln His Leu Ser Leu 475 470

Val Asn Leu Glu Pro Arg Ser Thr Tyr Arg Ile Cys Leu Val Pro Leu 485 490 495

Asp Ala Phe Asn Tyr Arg Ala Val Glu Asp Thr Ile Cys Ser Glu Ala 500 505 510

Thr Thr His Ala Ser Tyr Leu Asn Asn Gly Ser Asn Thr Ala Ser Ser 515 520 525

His Glu Gln Thr Thr Ser His Ser Met Gly Ser Pro Phe Leu Leu Ala 530 535 540

Gly Leu Ile Gly Gly Ala Val Ile Phe Val Leu Val Val Leu Leu Ser 545 550 555 560

Val Phe Cys Trp His Met His Lys Lys Gly Arg Tyr Thr Ser Gln Lys 565 570 575

Trp Lys Tyr Asn Arg Gly Arg Arg Lys Asp Asp Tyr Cys Glu Ala Gly
580 585 590

Thr Lys Lys Asp Asn Ser Ile Leu Glu Met Thr Glu Thr Ser Phe Gln 595 600 605

Ile Val Ser Leu Asn Asn Asp Gln Leu Leu Lys Gly Asp Phe Arg Leu 610 615 620

Gln Pro Ile Tyr Thr Pro Asn Gly Gly Ile Asn Tyr Thr Asp Cys His 625 630 635 640

Ile Pro Asn Asn Met Arg Tyr Cys Asn Ser Ser Val Pro Asp Leu Glu 645 650 655

His Cys His Thr

<210> 59

<211> 674

<212> PRT

<213> Homo sapiens

<400> 59

Met Val Val Ala His Pro Thr Ala Thr Ala Thr Thr Thr Pro Thr Ala
1 5 10 15

Thr Val Thr Ala Thr Val Val Met Thr Thr Ala Thr Met Asp Leu Arg

Asp Trp Leu Phe Leu Cys Tyr Gly Leu Ile Ala Phe Leu Thr Glu Val

Ile Asp Ser Thr Thr Cys Pro Ser Val Cys Arg Cys Asp Asn Gly Phe 50 60

Ile Tyr Cys Asn Asp Arg Gly Leu Thr Ser Ile Pro Ala Asp Ile Pro

Asp	Asp	Ala	Thr	Thr 85	Leu	Tyr	Leu	Gln	Asn 90	Asn	Gln	Ile	Asn	Asn 95	Ala
Gly	Ile	Pro	Gln 100	Asp	Leu	Lys	Thr	Lys 105	Val	Asn	Val	Gln	Val 110	Ile	Tyr
Leu	Tyr	Glu 115	Asn	Asp	Leu	Asp	Glu 120	Phe	Pro	Ile	Asn	Leu 125	Pro	Arg	Ser
Leu	Arg 130	Glu	Leu	His	Leu	Gln 135	Asp	Asn	Asn	Val	Arg 140	Thr	Ile	Ala	Arg
Asp 145	Ser	Leu	Ala	Arg	Ile 150	Pro	Leu	Leu	Glu	Lys 155	Leu	His	Leu	Asp	Asp 160
				165					170				Ala	175	
_			180					185					Ser 190		
		195					200					205	Asp		
	210					215					220		Asn		
225					230					235			Arg		240
				245					250				Ser	255	
			260					265					Ala 270		
		275					280					285	Pro		
	290					295					300		Ser		
305					310					315			Gly		320
				325					330				Cys	335	
			340					345					Val 350		
		355					360					365	Met		
Lys	Asp 370	Ile	Thr	Ser	Glu	Met 375	Asp	Glu	Cys	Phe	Glu 380	Thr	Gly	Pro	GIN

Gly Gly Val Ala Asn Ala Ala Ala Lys Thr Thr Ala Ser Asn His Ala Ser Ala Thr Thr Pro Gln Gly Ser Leu Phe Thr Leu Lys Ala Lys Arg 410 Pro Gly Leu Arg Leu Pro Asp Ser Asn Ile Asp Tyr Pro Met Ala Thr 425 Gly Asp Gly Ala Lys Thr Leu Ala Ile His Val Lys Ala Leu Thr Ala 440 Asp Ser Ile Arg Ile Thr Trp Lys Ala Thr Leu Pro Ala Ser Ser Phe 455 Arg Leu Ser Trp Leu Arg Leu Gly His Ser Pro Ala Val Gly Ser Ile Thr Glu Thr Leu Val Gln Gly Asp Lys Thr Glu Tyr Leu Leu Thr Ala 490 Leu Glu Pro Lys Ser Thr Tyr Ile Ile Cys Met Val Thr Met Glu Thr Ser Asn Ala Tyr Val Ala Asp Glu Thr Pro Val Cys Ala Lys Ala Glu 520 Thr Ala Asp Ser Tyr Gly Pro Thr Thr Thr Leu Asn Gln Glu Gln Asn Ala Gly Pro Met Ala Ser Leu Pro Leu Ala Gly Ile Ile Gly Gly Ala 555 550 Val Ala Leu Val Phe Leu Phe Leu Val Leu Gly Ala Ile Cys Trp Tyr Val His Gln Ala Gly Glu Leu Leu Thr Arg Glu Arg Ala Tyr Asn Arg 585 Gly Ser Arg Glu Lys Asp Asp Tyr Met Glu Ser Gly Thr Lys Lys Asp Asn Ser Ile Leu Glu Ile Arg Gly Pro Gly Leu Gln Met Leu Pro Ile Asn Pro Tyr Arg Ala Lys Glu Glu Tyr Val Val His Thr Ile Phe Pro 630 Ser Asn Gly Ser Ser Leu Cys Lys Ala Thr His Thr Ile Gly Tyr Gly Thr Thr Arg Gly Tyr Arg Asp Gly Gly Ile Pro Asp Ile Asp Tyr Ser 665 670

Tyr Thr

<210> 60 <211> 674 <212> PRT <213> Homo sapiens

<400> 60

Met Val Val Ala His Pro Thr Ala Thr Ala Thr Thr Thr Pro Thr Ala

1 5 10 15

Thr Val Thr Ala Thr Val Val Met Thr Thr Ala Thr Met Asp Leu Arg 20 25 30

Asp Trp Leu Phe Leu Cys Tyr Gly Leu Ile Ala Phe Leu Thr Glu Val 35 40 45

Ile Asp Ser Thr Thr Cys Pro Ser Val Cys Arg Cys Asp Asn Gly Phe
50 55 60

Ile Tyr Cys Asn Asp Arg Gly Leu Thr Ser Ile Pro Ala Asp Ile Pro 65 70 75 80

Asp Asp Ala Thr Thr Leu Tyr Leu Gln Asn Asn Gln Ile Asn Asn Ala 85 90 95

Gly Ile Pro Gln Asp Leu Lys Thr Lys Val Asn Val Gln Val Ile Tyr
100 105 110

Leu Tyr Glu Asn Asp Leu Asp Glu Phe Pro Ile Asn Leu Pro Arg Ser 115 120 125

Leu Arg Glu Leu His Leu Gln Asp Asn Asn Val Arg Thr Ile Ala Arg 130 135 140

Asp Ser Leu Ala Arg Ile Pro Leu Leu Glu Lys Leu His Leu Asp Asp 145 150 155 160

Asn Ser Val Ser Thr Val Ser Ile Glu Glu Asp Ala Phe Ala Asp Ser 165 170 175

Lys Gln Leu Lys Leu Leu Phe Leu Ser Arg Asn His Leu Ser Ser Ile 180 185 190

Pro Ser Gly Leu Pro His Thr Leu Glu Glu Leu Arg Leu Asp Asp Asn 195 200 205

Arg Ile Ser Thr Ile Pro Leu His Ala Phe Lys Gly Leu Asn Ser Leu 210 215 220

Arg Arg Leu Val Leu Asp Gly Asn Leu Leu Ala Asn Gln Arg Ile Ala 225 230 235 240

Asp Asp Thr Phe Ser Arg Leu Gln Asn Leu Thr Glu Leu Ser Leu Val 245 250 255

Arg Asn Ser Leu Ala Ala Pro Pro Leu Asn Leu Pro Ser Ala His Leu

			260					265					270		
		_		_	~1.				T1 -	<b>0</b>	TT -	T10		T	N a n
Gln	Lys	Leu 275	Tyr	Leu	GIn	Asp	280	Ala	11e	ser	HIS	285	PIO	ıyı	ASII
Thr	Leu 290	Ala	Lys	Met	Arg	Glu 295	Leu	Glu	Arg	Leu	Asp 300	Leu	Ser	Asn	Asn
Asn 305	Leu	Thr	Thr	Leu	Pro 310	Arg	Gly	Leu	Phe	Asp 315	Asp	Leu	Gly	Asn	Leu 320
Ala	Gln	Leu	Leu	Leu 325	Arg	Asn	Asn	Pro	Trp 330	Phe	Cys	Gly	Cys	Asn 335	Leu
Met	Trp	Leu	Arg 340	Asp	Trp	Val	Lys	Ala 345	Arg	Ala	Ala	Val	Val 350	Asn	Val
Arg	Gly	Leu 355	Met	Cys	Gln	Gly	Pro 360	Glu	Lys	Val	Arg	Gly 365	Met	Ala	Ile
Lys	Asp 370	Ile	Thr	Ser	Glu	Met 375	Asp	Glu	Cys	Phe	Glu 380	Thr	Gly	Pro	Gln
Gly 385	Gly	Val	Ala	Asn	Ala 390	Ala	Ala	Lys	Thr	Thr 395	Ala	Ser	Asn	His	Ala 400
Ser	Ala	Thr	Thr	Pro 405	Gln	Gly	Ser	Leu	Phe 410	Thr	Leu	Lys	Ala	Lys 415	Arg
Pro	Gly	Leu	Arg 420	Leu	Pro	Asp	Ser	Asn 425	Ile	Asp	Tyr	Pro	Met 430	Ala	Thr
Gly	Asp	Gly 435	Ala	Lys	Thr	Leu	Ala 440	Ile	His	Val	Lys	Ala 445	Leu	Thr	Ala
Asp	Ser 450	Ile	Arg	Ile	Thr	Trp 455	Lys	Ala	Thr	Leu	Pro 460	Ala	Ser	Ser	Phe
Arg 465	Leu	Ser	Trp	Leu	Arg 470	Leu	Gly	His	Ser	Pro 475	Ala	Val	Gly	Ser	Ile 480
Thr	Glu	Thr	Leu	Val 485	Gln	Gly	Asp	Lys	Thr 490	Glu	Tyr	Leu	Leu	Thr 495	Ala
Leu	Glu	Pro	Lys 500	Ser	Thr	Tyr	Ile	Ile 505	Cys	Met	Val	Thr	Met 510	Glu	Thr
Ser	Asn	Ala 515	Tyr	Val	Ala	Asp	Glu 520	Thr	Pro	Val	Cys	Ala 525	Lys	Ala	Glu
Thr	Ala 530	Asp	Ser	Tyr	Gly	Pro 535	Thr	Thr	Thr	Leu	Asn 540	Gln	Glu	Gln	Asn

Ala Gly Pro Met Ala Ser Leu Pro Leu Ala Gly Ile Ile Gly Gly Ala

Val Ala Leu Val Phe Leu Phe Leu Val Leu Gly Ala Ile Cys Trp Tyr

565	5 <b>7</b> 0	575

Val His Gln Ala Gly Glu Leu Leu Thr Arg Glu Arg Ala Tyr Asn Arg 580 585 590

Gly Ser Arg Glu Lys Asp Asp Tyr Met Glu Ser Gly Thr Lys Lys Asp 595 600 605

Asn Ser Ile Leu Glu Ile Arg Gly Pro Gly Leu Gln Met Leu Pro Ile 610 615 620

Asn Pro Tyr Arg Ala Lys Glu Glu Tyr Val Val His Thr Ile Phe Pro 625 630 635 640

Ser Asn Gly Ser Ser Leu Cys Lys Ala Thr His Thr Ile Gly Tyr Gly
645 650 655

Thr Thr Arg Gly Tyr Arg Asp Gly Gly Ile Pro Asp Ile Asp Tyr Ser
660 665 670

Tyr Thr

<210> 61

<211> 246

<212> PRT

<213> Homo sapiens

<400> 61

Pro Met Ala Thr Gly Asp Gly Ala Lys Thr Leu Ala Ile His Val Lys

1 10 15

Ala Leu Thr Ala Asp Ser Ile Arg Ile Thr Trp Lys Ala Thr Leu Pro 20 25 30

Ala Ser Ser Phe Arg Leu Ser Trp Leu Arg Leu Gly His Ser Pro Ala 35 40 45

Val Gly Ser Ile Thr Glu Thr Leu Val Gln Gly Asp Lys Thr Glu Tyr 50 55 60

Leu Leu Thr Ala Leu Glu Pro Lys Ser Thr Tyr Ile Ile Cys Met Val 65 70 75 80

Thr Met Glu Thr Ser Asn Ala Tyr Val Ala Asp Glu Thr Pro Val Cys
85 90 95

Ala Lys Ala Glu Thr Ala Asp Ser Tyr Gly Pro Thr Thr Leu Asn
100 105 110

Gln Glu Gln Asn Ala Gly Pro Met Ala Ser Leu Pro Leu Ala Gly Ile 115 120 125

Ile Gly Gly Ala Val Ala Leu Val Phe Leu Phe Leu Val Leu Gly Ala 130 140 Ile Cys Trp Tyr Val His Gln Ala Gly Glu Leu Leu Thr Arg Glu Arg 145 150 155 160

Ala Tyr Asn Arg Gly Ser Arg Lys Lys Asp Asp Tyr Met Glu Ser Gly
165 170 175

Thr Lys Lys Asp Asn Ser Ile Leu Glu Ile Arg Gly Pro Gly Leu Gln
180 185 190

Met Leu Pro Ile Asn Pro Tyr Arg Ala Lys Glu Glu Tyr Val Val His 195 200 205

Thr Ile Phe Pro Ser Asn Gly Ser Ser Leu Cys Lys Ala Thr His Thr 210 215 220

Ile Gly Tyr Gly Thr Thr Arg Gly Tyr Arg Asp Gly Gly Ile Pro Asp 225 230 235 240

Ile Asp Tyr Ser Tyr Thr

210> 62

<211> 378

<212> PRT

<213> Homo sapiens

<400> 62

Ile Ser Asn Asn Gln Leu Arg Met Leu Thr Gln Gly Val Phe Asp Asn 1 5 10 15

Leu Ser Asn Leu Lys Gln Leu Thr Ala Arg Asn Asn Pro Trp Phe Cys
20 25 30

Asp Cys Ser Ile Lys Trp Val Thr Glu Trp Leu Lys Tyr Ile Pro Ser 35 40 45

Ser Leu Asn Val Arg Gly Phe Met Cys Gln Gly Pro Glu Gln Val Arg 50 55 60

Gly Met Ala Val Arg Glu Leu Asn Met Asn Leu Leu Ser Cys Pro Thr 65 70 75 80

Thr Thr Pro Gly Leu Pro Leu Phe Thr Pro Ala Pro Ser Thr Ala Ser 85 90 95

Pro Thr Thr Gln Pro Pro Thr Leu Ser Ile Pro Asn Pro Ser Arg Ser 100 105 110

Tyr Thr Pro Pro Thr Pro Thr Thr Ser Lys Leu Pro Thr Ile Pro Asp 115 120 125

Trp Asp Gly Arg Glu Arg Val Thr Pro Pro Ile Ser Glu Arg Ile Gln
130 135 140

Leu Ser Ile His Phe Val Asn Asp Thr Ser Ile Gln Val Ser Trp Leu 145 150 155 160 Ser Leu Phe Thr Val Met Ala Tyr Lys Leu Thr Trp Val Lys Met Gly His Ser Leu Val Gly Gly Ile Val Gln Glu Arg Ile Val Ser Gly Glu 185 Lys Gln His Leu Ser Leu Val Asn Leu Glu Pro Arg Ser Thr Tyr Arg 200 Ile Cys Leu Val Pro Leu Asp Ala Phe Asn Tyr Arg Ala Val Glu Asp 215 Thr Ile Cys Ser Glu Ala Thr Thr His Ala Ser Tyr Leu Asn Asn Gly 235 230 Ser Asn Thr Ala Ser Ser His Glu Gln Thr Thr Ser His Ser Met Gly 250 Ser Pro Phe Leu Leu Ala Gly Leu Ile Gly Gly Ala Val Ile Phe Val Leu Val Val Leu Leu Ser Val Phe Cys Trp His Met His Lys Lys Gly Arg Tyr Thr Ser Gln Lys Trp Lys Tyr Asn Arg Gly Arg Arg Lys Asp 295 Asp Tyr Cys Glu Ala Gly Thr Lys Lys Asp Asn Ser Ile Leu Glu Met Thr Glu Thr Ser Phe Gln Ile Val Ser Leu Asn Asn Asp Gln Leu Leu 330 Lys Gly Asp Phe Arg Leu Gln Pro Ile Tyr Thr Pro Asn Gly Gly Ile Asn Tyr Thr Asp Cys His Ile Pro Asn Asn Met Arg Tyr Cys Asn Ser Ser Val Pro Asp Leu Glu His Cys His Thr 370 <210> 63 <211> 338 <212> PRT <213> Gallus gallus <400> 63 Val His Ser Val Trp Thr Arg Thr Val Arg Gln Val Tyr Asn Glu Leu

Asp Pro Glu His Trp Ser His Tyr Thr Phe Glu Cys Pro Gln Glu Cys

Phe Cys Pro Pro Ser Phe Pro Asn Ala Leu Tyr Cys Asp Asn Lys Gly

40 45

35

Val Ile

Leu	Lys 50	Glu	Ile	Pro	Ala	Ile 55	Pro	Ala	Arg	Ile	Trp 60	Tyr	Leu	Tyr	Leu
Gln 65	Asn	Asn	Leu	Ile	Glu 70	Thr	Ile	Ser	Glu	Lys <b>7</b> 5	Pro	Phe	Val	Asn	Ala 80
Thr	His	Leu	Arg	Trp 85	Ile	Asn	Leu	Asn	Lys 90	Asn	Lys	Ile	Thr	Asn 95	Asn
Gly	Ile	Glu	Ser 100	Gly	Val	Leu	Ser	Lys 105	Leu	Lys	Arg	Leu	Leu 110	Tyr	Leu
Phe	Leu	Glu 115	Asp	Asn	Glu	Leu	Glu 120	Glu	Val	Pro	Ala	Pro 125	Leu	Pro	Val
Gly	Leu 130	Glu	Gln	Leu	Arg	Leu 135	Ala	Arg	Asn	Lys	Ile 140	Ser	Arg	Ile	Pro
Glu 145	Gly	Val	Phe	Ser	Asn 150	Leu	Glu	Asn	Leu	Thr 155	Met	Leu	Asp	Leu	His 160
Gln	Asn	Asn	Leu	Leu 165	Asp	Ser	Ala	Leu	Gln 170	Ser	Asp	Thr	Phe	Gln 175	Gly
Leu	Asn	Ser	Leu 180	Met	Gln	Leu	Asn	Ile 185	Ala	Lys	Asn	Ser	Leu 190	Lys	Lys
Met	Pro	Leu 195	Ser	Ile	Pro	Ala	Asn 200	Thr	Leu	Gln	Leu	Phe 205	Leu	Asp	Asn
Asn	Ser 210	Ile	Glu	Val	Ile	Pro 215	Glu	Asn	Tyr	Phe	Ser 220	Ala	Ile	Pro	Lys
Val 225	Thr	Phe	Leu	Arg	Leu 230	Asn	Tyr	Asn	Lys	Leu 235	Ser	Asp	Asp	Gly	Ile 240
Pro	Pro	Asn	Gly	Phe 245	Asn	Val	Ser	Ser	Ile 250	Leu	Asp	Leu	Gln	Leu 255	Ser
His	Asn	Gln	Leu 260	Thr	Lys	Ile	Pro	Pro 265	Ile	Asn	Ala	His	Leu 270	Glu	His
Leu	His	Leu 275	Asp	His	Asn	Arg	11e 280	Lys	Ser	Val	Asn	Gly 285	Thr	Gln	Ile
Cys	Pro 290	Val	Ser	Ile	Ala	Val 295	Ala	Glu	Asp	Tyr	Gly 300	Leu	Tyr	Gly	Asn
Ile 305	Pro	Arg	Leu	Arg	Tyr 310	Leu	Arg	Leu	Asp	Gly 315	Asn	Glu	Ile	Gln	Pro 320
Pro	Ile	Pro	Leu	Asp 325	Ile	Met	Ile	Суѕ	Phe 330	Gln	Leu	Leu	Gln	Ala 335	Val

<210> 64 <211> 326 <212> PRT <213> Bos taurus

<400> 64

Pro Tyr Glu Pro Tyr Pro Thr Gly Glu Glu Gly Pro Ala Tyr Ala Tyr 1 5 10 15

Gly Ser Pro Pro Gln Pro Glu Pro Arg Asp Cys Pro Gln Glu Cys Asp
20 25 30

Cys Pro Pro Asn Phe Pro Thr Ala Met Tyr Cys Asp Asn Arg Asn Leu 35 40  $\cdot$  45

Lys Tyr Leu Pro Phe Val Pro Ser Arg Met Lys Tyr Val Tyr Phe Gln 50 60

Asn Asn Gln Ile Ser Ser Ile Gln Glu Gly Val Phe Asp Asn Ala Thr 65 70 75 80

Gly Leu Leu Trp Ile Ala Leu His Gly Asn Gln Ile Thr Ser Asp Lys
85 90 95

Val Gly Lys Lys Val Phe Ser Lys Leu Arg His Leu Glu Arg Leu Tyr 100 105 110

Leu Asp His Asn His Leu Thr Arg Ile Pro Ser Pro Leu Pro Arg Ser 115 120 125

Leu Arg Glu Leu His Leu Asp His Asn Gln Ile Ser Arg Val Pro Asn 130 135 140

Asn Ala Leu Glu Gly Leu Glu Asn Leu Thr Ala Leu Tyr Leu His His 145 150 155 160

Glu Ile Gln Glu Val Gly Ser Ser Met Lys Gly Leu Arg Ser Leu Ile 165 170 175

Leu Leu Asp Leu Ser Tyr Asn His Leu Arg Lys Val Pro Asp Gly Leu 180 185 190

Pro Ser Ala Leu Glu Gln Leu Tyr Leu Glu His Asn Asn Val Phe Ser 195 200 205

Val Pro Asp Ser Tyr Phe Arg Gly Ser Pro Lys Leu Leu Tyr Val Arg 210 215 220

Leu Ser His Asn Ser Leu Thr Asn Asn Gly Leu Ala Ser Asn Thr Phe 225 230 235 240

Asn Ser Ser Ser Leu Leu Glu Leu Asp Leu Ser Tyr Asn Gln Leu Gln
245 250 255

```
Lys Ile Pro Pro Val Ser Thr Asn Leu Glu Asn Leu Tyr Leu Gln Gly
                                 265
Asn Arg Ile Asn Glu Phe Ser Ile Ser Ser Phe Cys Thr Val Val Asp
        275
                            280
                                                 285
Val Met Asn Phe Ser Lys Leu Gln Val Gln Arg Leu Asp Gly Asn Glu
                        295
Ile Lys Arg Ser Ala Met Pro Ala Asp Ala Pro Leu Cys Leu Arg Leu
305
Ala Ser Leu Ile Glu Ile
                325
<210> 65
<211> 1020
<212> DNA
<213> Homo sapiens
<400> 65
gegegeggeg aagtgaattt getggaeaeg tegaecatee aeggggaetg gggetggete 60
acgtatccgg ctcatgggtg ggactccatc aacgaggtgg acgagtcctt ccaqcccatc 120
cacacgtacc aggtttqcaa cqtcatqaqc cccaaccaqa acaactqqct qcqcacqaqc 180
tgggtccccc gagacggcqc ccgqcqcqtc tatqctqaga tcaaqtttac cctqcqcqac 240
tgcaacagca tgcctggtgt gctgggcacc tgcaaggaga ccttcaacct ctactacctg 300
gagteggace gegacetggg ggeeageaca caagaaagee agtteeteaa aategacace 360
attgeggeeg acqaqaqett cacaqqtqee qacettqqtq tqeqqeqtet caaqetcaac 420
acggaggtgc gcagtgtggg tcccctcagc aagcgcggct tctacctggc cttccaggac 480
ataggtgcct gcctggccat cctctctc cgcatctact ataagaagtg ccctgccatg 540
gtgcgcaatc tggctgcctt ctcggaggca gtgacggggg ccgactcgtc ctcactggtg 600
gaggtgaggg gccagtgcgt gcggcactca gaggagcggg acacacccaa gatgtactgc 660
agegeggagg gegagtgget egtgeecate ggeaaatgeg tgtgeagtge eggetaegag 720
gageggeggg atgcctgtgt ggcctgtgag ctgggcttct acaagtcagc ccctggggac 780
cagetgtgtg ceegetgeee tecceacage cacteegeag etccageege ceaageetge 840
cactgtgacc tcagctacta ccgtgcagcc ctggacccgc cgtcctcagc ctgcacccgg 900
ccaccetegg caccagtgaa cetgatetee agtgtgaatg ggacateagt gaetetggag 960
tgggcccctc ccctggaccc aggtggccgc agtgacatca cctacaatgc cqtqtqccqc 1020
<210> 66
<211> 515
<212> PRT
<213> Homo sapiens
Ala Arq Gly Glu Val Asn Leu Leu Asp Thr Ser Thr Ile His Gly Asp
Trp Gly Trp Leu Thr Tyr Pro Ala His Gly Trp Asp Ser Ile Asn Glu
                                 25
Val Asp Glu Ser Phe Gln Pro Ile His Thr Tyr Gln Val Cys Asn Val
```

Asp Gly Ala Arg Arg Val Tyr Ala Glu Ile Lys Phe Thr Leu Arg Asp 75 Cys Asn Ser Met Pro Gly Val Leu Gly Thr Cys Lys Glu Thr Phe Asn Leu Tyr Tyr Leu Glu Ser Asp Arg Asp Leu Gly Ala Ser Thr Gln Glu 100 105 Ser Gln Phe Leu Lys Ile Asp Thr Ile Ala Ala Asp Glu Ser Phe Thr Gly Ala Asp Leu Gly Val Arg Arg Leu Lys Leu Asn Thr Glu Val Arg Ser Val Gly Pro Leu Ser Lys Arg Gly Phe Tyr Leu Ala Phe Gln Asp Ile Gly Ala Cys Leu Ala Ile Leu Ser Leu Arg Ile Tyr Tyr Lys Lys 170 Cys Pro Ala Met Val Arg Asn Leu Ala Ala Phe Ser Glu Ala Val Thr Gly Ala Asp Ser Ser Ser Leu Val Glu Val Arg Gly Gln Cys Val Arg His Ser Glu Glu Arg Asp Thr Pro Lys Met Tyr Cys Ser Ala Glu Gly Glu Trp Leu Val Pro Ile Gly Lys Cys Val Cys Ser Ala Gly Tyr Glu 230 235 Glu Arg Arg Asp Ala Cys Val Ala Cys Glu Leu Gly Phe Tyr Lys Ser 245 Ala Pro Gly Asp Gln Leu Cys Ala Arg Cys Pro Pro His Ser His Ser 265 Ala Ala Pro Ala Ala Gln Ala Cys His Cys Asp Leu Ser Tyr Tyr Arg 275

Met Ser Pro Asn Gln Asn Asn Trp Leu Arg Thr Ser Trp Val Pro Arg

Trp Ala Pro Pro Leu Asp Pro Gly Gly Arg Ser Asp Ile Thr Tyr Asn 325 330 335

Ala Ala Leu Asp Pro Pro Ser Ser Ala Cys Thr Arg Pro Pro Ser Ala

Pro Val Asn Leu Ile Ser Ser Val Asn Gly Thr Ser Val Thr Leu Glu

295

305

Ala Val Cys Arg Arg Cys Pro Trp Ala Leu Ser Arg Cys Glu Ala Cys 340 345 350

315

Gly Ser Gly Thr Arg Phe Val Pro Gln Gln Thr Ser Leu Val Gln Ala Ser Leu Leu Val Ala Asn Leu Leu Ala His Met Asn Tyr Ser Phe Trp 380 370 375 Ile Glu Ala Val Asn Gly Val Ser Asp Leu Ser Pro Glu Pro Arg Arg 390 395 Ala Ala Val Val Asn Ile Thr Thr Asn Gln Ala Ala Pro Ser Gln Val Val Val Ile Arg Gln Glu Arg Ala Gly Gln Thr Ser Val Ser Leu Leu Trp Gln Glu Pro Glu Gln Pro Asn Gly Ile Ile Leu Glu Tyr Glu Ile Lys Tyr Tyr Glu Lys Asp Lys Glu Met Gln Ser Tyr Ser Thr Leu Lys 455 Ala Val Thr Thr Arg Ala Thr Val Ser Gly Leu Lys Pro Gly Thr Arg 470 475 Tyr Val Phe Gln Val Arg Ala Arg Thr Ser Ala Gly Cys Gly Arg Phe Ser Gln Ala Met Glu Val Glu Thr Gly Lys Pro Arg Pro Arg Tyr Asp 505 Thr Arg Thr 515 <210> 67 <211> 1992 <212> DNA <213> Homo sapiens <400> 67 atgqtqqtqq cacaccccac cgccactgcc accaccacgc ccactgccac tgtcacggcc 60 accqttgtga tgaccacggc caccatggac ctgcgggact ggctgttcct ctgctacggg 120 ctcatcqcct tcctqacqqa qqtcatcqac aqcaccacct gcccctcggt gtgccgctgc 180 qacaacqqct tcatctactq caacqaccqq qqactcacat ccatccccqc agatatccct 240 qatqatqcca ccaccctcta cctgcagaac aaccagatca acaacgccgg catcccccag 300 qacctcaaqa ccaaqqtcaa cqtqcaqqtc atctacctat acgagaatga cctggatgag 360 ttccccatca acctgccccg ctccctccgg gagctgcacc tgcaggacaa caatgtgcgc 420 accattgeca gggacteget ggcccgcatc cegetgetgg agaagetgea cetggatgac 480 aactccgtgt ccaccgtcag cattgaggag gacgccttcg ccgacagcaa acagctcaag 540 ctgctcttcc tgagccggaa ccacctgagc agcatcccct cggggctgcc gcacacgctg 600 qaqqaqctqc qqctqqatqa caaccqcatc tccaccatcc cgctqcatgc cttcaagggc 660 ctcaacagcc tgcggcgcct ggtgctggac ggtaacctgc tggccaacca gcgcatcgcc 720 qacqacacct tcagccgcct acagaacctc acagagctct cgctggtgcg caattcgctg 780 qccqcqccac ccctctacct qcaggacaat qccatcagcc acatccccta caacacgctg 840 gccaagatgc gtgagctgga gcggctggac ctgtccaaca acaacctgac cacgctgccc 900 cqcqqcctqt tcqacqacct qqqqaacctq qcccaqctqc tgctcaggaa caacccttgg 960

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<213> Homo sapiens

<400> 68

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Asp Trp Leu Phe Leu Cys Tyr Gly Leu Ile Ala Phe Leu Thr Glu Val 35 40 45

Ile Asp Ser Thr Thr Cys Pro Ser Val Cys Arg Cys Asp Asn Gly Phe 50 60

Ile Tyr Cys Asn Asp Arg Gly Leu Thr Ser Ile Pro Ala Asp Ile Pro 65 70 75 80

Asp Asp Ala Thr Thr Leu Tyr Leu Gln Asn Asn Gln Ile Asn Asn Ala 85 90 95

Gly Ile Pro Gln Asp Leu Lys Thr Lys Val Asn Val Gln Val Ile Tyr 100 105 110

Leu Tyr Glu Asn Asp Leu Asp Glu Phe Pro Ile Asn Leu Pro Arg Ser 115 120 125

Leu Arg Glu Leu His Leu Gln Asp Asn Asn Val Arg Thr Ile Ala Arg 130 140

Asp Ser Leu Ala Arg Ile Pro Leu Leu Glu Lys Leu His Leu Asp Asp 145 150 155 160

Asn Ser Val Ser Thr Val Ser Ile Glu Glu Asp Ala Phe Ala Asp Ser 165 170 175 Lys Gln Leu Lys Leu Leu Phe Leu Ser Arg Asn His Leu Ser Ser Ile 180 185 Pro Ser Gly Leu Pro His Thr Leu Glu Glu Leu Arg Leu Asp Asp Asn Arg Ile Ser Thr Ile Pro Leu His Ala Phe Lys Gly Leu Asn Ser Leu 215 Arg Arg Leu Val Leu Asp Gly Asn Leu Leu Ala Asn Gln Arg Ile Ala 230 Asp Asp Thr Phe Ser Arg Leu Gln Asn Leu Thr Glu Leu Ser Leu Val 250 Arg Asn Ser Leu Ala Ala Pro Pro Leu Tyr Leu Gln Asp Asn Ala Ile Ser His Ile Pro Tyr Asn Thr Leu Ala Lys Met Arg Glu Leu Glu Arg 280 Leu Asp Leu Ser Asn Asn Asn Leu Thr Thr Leu Pro Arg Gly Leu Phe 295 Asp Asp Leu Gly Asn Leu Ala Gln Leu Leu Leu Arg Asn Asn Pro Trp 310 315 Phe Cys Gly Cys Asn Leu Met Trp Leu Arg Asp Trp Val Lys Ala Arg 325 330 Ala Ala Val Val Asn Val Arg Gly Leu Met Cys Gln Gly Pro Glu Lys 345 Val Arg Gly Met Ala Ile Lys Asp Ile Thr Ser Glu Val Glu Ser Val Leu Arg Arg Ala Pro Gln Gly Gly Val Ala Asn Ala Ala Ala Lys Thr Thr Ala Ser Asn His Ala Ser Ala Thr Thr Pro Gln Gly Ser Leu Phe 385 Thr Leu Lys Ala Lys Arg Pro Gly Leu Arg Leu Pro Asp Ser Asn Ile 410 Asp Tyr Pro Met Ala Thr Gly Asp Gly Ala Lys Thr Leu Ala Ile His 425 Val Lys Ala Leu Thr Ala Asp Ser Ile Arg Ile Thr Trp Lys Ala Thr Leu Pro Ala Ser Ser Phe Arg Leu Ser Trp Leu Arg Leu Gly His Ser 455 450 Pro Ala Val Gly Ser Ile Thr Glu Thr Leu Val Gln Gly Asp Lys Thr 470 475 465

Glu Tyr Leu Leu Thr Ala Leu Glu Pro Lys Ser Thr Tyr Ile Ile Cys Met Val Thr Met Glu Thr Ser Asn Ala Tyr Val Ala Asp Glu Thr Pro 505 Val Cys Ala Lys Ala Glu Thr Ala Asp Ser Tyr Gly Pro Thr Thr 515 520 525 Leu Asn Gln Glu Gln Asn Ala Gly Pro Met Ala Ser Leu Pro Leu Ala 535 Gly Ile Ile Gly Gly Ala Val Ala Leu Val Phe Leu Phe Leu Val Leu 545 Gly Ala Ile Cys Trp Tyr Val His Gln Ala Gly Glu Leu Leu Thr Arg Glu Arg Ala Tyr Asn Arg Gly Ser Arg Lys Lys Asp Asp Tyr Met Glu 585 Ser Gly Thr Lys Lys Asp Asn Ser Ile Leu Glu Ile Arg Gly Pro Gly 600 Leu Gln Met Leu Pro Ile Asn Pro Tyr Arg Ala Lys Glu Glu Tyr Val 615 Val His Thr Ile Phe Pro Ser Asn Gly Ser Ser Leu Cys Lys Ala Thr His Thr Ile Gly Tyr Gly Thr Thr Arg Gly Tyr Arg Asp Gly Gly Ile 650 Pro Asp Ile Asp Tyr Ser Tyr Thr 660 <210> 69 <211> 26 <212> DNA <213> Homo sapiens <400> 69 caacgtgcag gtcatctacc tatacg 26 <210> 70 <211> 25 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence:oligonucleotide primer

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<211> 54
<212> PRT
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Arg Glu Thr Asn Pro Arg Arg Leu Glu Asp Gln Glu Asp Leu Arg Cys

Ala Ser Pro Glu Ser Leu Arg Gly Gln Pro Leu Leu Glu Leu Leu Pro 40

Ser Asp Phe Ser Cys Pro

<210> 72 <211> 84 <212> PRT

<213> Artificial Sequence

<223> Description of Artificial Sequence:consensus sequence

<400> 72

Pro Ser Ala Pro Thr Asn Leu Thr Val Thr Asp Val Thr Ser Thr Ser

Leu Thr Leu Ser Trp Ser Pro Pro Thr Gly Asn Gly Pro Ile Thr Gly

Tyr Glu Val Thr Tyr Arg Gln Pro Lys Asn Gly Gly Glu Trp Asn Glu

Leu Thr Val Pro Gly Thr Thr Thr Ser Tyr Thr Leu Thr Gly Leu Lys 55

Pro Gly Thr Glu Tyr Glu Val Arg Val Gln Ala Val Asn Gly Gly 65 70 75

Gly Pro Glu Ser

<210> 73 <211> 23

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Gly Leu Phe Ser Asn Leu Pro
                             20
 <210> 74
 <211> 949
 <212> DNA
 <213> Homo sapiens
<400> 74
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acceptiging the transfer of the second secon
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gacaacggct tcatctactg caacgaccgg ggactcacat ccatccccgc agatatccct 240
gatgacgcca ccaccctcta tctgcagaac aaccagatca acaacgctgg catcccccag 300
gacctcaaga ccaaggtcaa cgtgcaggtc atctacctat acgagaatga cctggatgag 360
ttccccatca acctgccccg ctccctccqq qaqctqcacc tqcaqqacaa caatqtqcqc 420
accattgcca gggactcgct ggcccgcatc ccqctqctqq aqaaqctqca cctqqatqac 480
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ctgctcttcc tgagccggaa ccacctgagc agcatcccct cqqqqctqcc qcacacqctq 600
gaggagetge ggetggatga caacegeate tecaceatee egetgeatge etteaaggge 660
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gacgacacct teageegeet acagaacete acagagetet egetggtgeg caattegetg 780
gccgcgccac ccctcaacct gcccagcgcc cacctgcaga aactctacct gcaggacaat 840
gccatcagcc acatccccta caacacgctg gccaagatgc gtgagctgga gcggctggac 900
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<211> 674
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<213> Homo sapiens
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Thr Val Thr Ala Thr Val Val Met Thr Thr Ala Thr Met Asp Leu Arq
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Asp Trp Leu Phe Leu Cys Tyr Gly Leu Ile Ala Phe Leu Thr Glu Val
Ile Asp Ser Thr Thr Cys Pro Ser Val Cys Arg Cys Asp Asn Gly Phe
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Ile Tyr Cys Asn Asp Arg Gly Leu Thr Ser Ile Pro Ala Asp Ile Pro

Lys Asp Ile Thr Ser Glu Met Asp Glu Cys Phe Glu Thr Gly Pro Gln

	370					375					380				
Gly 385	Gly	Val	Ala	Asn	Ala 390	Ala	Ala	Lys	Thr	Thr 395	Ala	Ser	Asn	His	Ala 400
Ser	Ala	Thr	Thr	Pro 405	Gln	Gly	Ser	Leu	Phe 410	Thr	Leu	Lys	Ala	Lys 415	Arg
Pro	Gly	Leu	Arg 420	Leu	Pro	Asp	Ser	Asn 425	Ile	Asp	Tyr	Pro	Met 430	Ala	Thr
Gly	Asp	Gly 435	Ala	Lys	Thr	Leu	Ala 440	Ile	His	Val	Lys	Ala 445	Leu	Thr	Ala
Asp	Ser 450	Ile	Arg	Ile	Thr	Trp 455	Lys	Ala	Thr	Leu	Pro 460	Ala	Ser	Ser	Phe
Arg 465	Leu	Ser	Trp	Leu	Arg 470	Leu	Gly	His	Ser	Pro 475	Ala	Val	Gly	Ser	Ile 480
Thr	Glu	Thr	Leu	Val 485	Gln	Gly	Asp	Lys	Thr 490	Glu	Tyr	Leu	Leu	Thr 495	Ala
Leu	Glu	Pro	Lys 500	Ser	Thr	Tyr	Ile	Ile 505	Cys	Met	Val	Thr	Met 510	Glu	Thr
Ser	Asn	Ala 515	Tyr	Val	Ala	Asp	Glu 520	Thr	Pro	Val	Cys	Ala 525	Lys	Ala	Glu
Thr	Ala 530	Asp	Ser	Tyr	Gly	Pro 535	Thr	Thr	Thr	Leu	Asn 540	Gln	Glu	Gln	Asn
Ala 545	Gly	Pro	Met	Ala	Ser 550	Leu	Pro	Leu	Ala	Gly 555	Ile	Ile	Gly	Gly	Ala 560
Val	Ala	Leu	Val	Phe 565	Leu	.Phe	Leu	Val	Leu 570	Gly	Ala	Ile	Cys	Trp 575	Tyr
Val	His	Gln	Ala 580	Gly	Glu	Leu	Leu	Thr 585	Arg	Glu	Arg	Ala	Tyr 590	Asn	Arg
Gly	Ser	Arg 595	Glu	Lys	Asp	Asp	Tyr 600	Met	Glu	Ser	Gly	Thr 605	Lys	Lys	Asp
Asn	Ser 610	Ile	Leu	Glu	Ile	Arg 615	Gly	Pro	Gly	Leu	Gln 620	Met	Leu	Pro	Ile
Asn 625	Pro	Tyr	Arg	Ala	Lys 630	Glu	Glu	Tyr	Val	Val 635	His	Thr	Ile	Phe	Pro 640
Ser	Asn	Gly	Ser	Ser 645	Leu	Cys	Lys	Ala	Thr 650	His	Thr	Ile	Gly	Tyr 655	Gly
Thr	Thr	Arg	Gly 660	Tyr	Arg	Asp	Gly	Gly 665	Ile	Pro	Asp	Ile	Asp 670	Tyr	Ser
Tyr	Thr														

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<211> 31
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence:consensus
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Cys Ser Gly Arg Gly Leu Thr Leu Glu Val Pro Arg Asp Leu Pro
<210> 77
<211> 23
<212> PRT
<213> Artificial Sequence
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Gly Leu Phe Ser Asn Leu Pro
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<210> 78
<211> 23
<212> PRT
<213> Artificial Sequence
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<400> 78
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Gly Leu Phe Ser Asn Leu Pro
             20
<210> 79
<211> 23
<212> PRT
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<213> Artificial Sequence
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Gly Leu Phe Ser Asn Leu Pro
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Gly Leu Phe Ser Asn Leu Pro
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<211> 23
<212> PRT
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Gly Leu Phe Ser Asn Leu Pro
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<210> 82
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<212> PRT
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      sequence
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Asn Leu Glu Glu Leu Asp Leu Ser Asn Asn Leu Thr Ser Leu Pro Pro 5 10 Gly Leu Phe Ser Asn Leu Pro 20 <210> 83 <211> 23 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence:consensus sequence <400> 83 Asn Leu Glu Glu Leu Asp Leu Ser Asn Asn Leu Thr Ser Leu Pro Pro 10 Gly Leu Phe Ser Asn Leu Pro 20 <210> 84 <211> 23 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence:consensus sequence <400> 84 Asn Leu Glu Glu Leu Asp Leu Ser Asn Asn Leu Thr Ser Leu Pro Pro 10 Gly Leu Phe Ser Asn Leu Pro 20